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Heparin-II binding domain of fibronectin having an amino acid sequence represented by the formula:

Ala lle Pro Ala Pro Thr Asr Leu Lys Phe Thr Gln Val Thr Pro Thr Ser Leu Ser Ala Gln Trp Thr Pro Pro Asn Val Gln Leu Thr Gly Tyr Arg Val Arg Val Thr Pro Lys Glu Lys Thr Gly Pro Met Lys Glu lle Asn Leu Ala Pro Asp Ser Ser Ser Val Val Val Ser Gly Leu Met Val Ala Thr Lys Tyr Glu Val Ser Val Tyr Ala Leu Lys Asp Thr Leu Thr Ser Arg Pro Ala Gln Gly Val Val Thr Thr Leu Glu Asn Val Ser Pro Pro Arg Arg Ala Arg Val Thr Asp Ala Thr Glu Thr Thr lle Thr lle Ser Trp Arg Thr Lys Thr Glu Thr lle Thr Gly Phe Gln Val Asp Ala Val Pro Ala Asn Gly Gln Thr Pro lle Gln Arg Thr lle Lys Pro Asp Val Arg Ser Tyr Thr lle Thr Gly Leu Gln Pro Gly Thr Asp Tyr Lys lle Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val Val lle Asp Ala Ser Thr Ala lle Asp Ala Pro Ser Asn Leu Arg Phe Leu Ala Thr Thr Pro Asn Ser Leu Leu Val Ser Trp Gln Pro Pro Arg Ala Arg lle Thr Gly Tyr lle lle Lys Tyr Glu Lys Pro Gly Ser Pro/Pro Arg Glu Val Val Pro Arg Pro Arg Pro Gly Val Thr Glu Ala Thr Ile Thr Gly Leu Glu Pro Leu Ile Gly Arg Lys Lys Thr

or a sufficiently similar amino acid sequence thereto to exhibit the ability to bind retroviruses.

93. The cellular population of claim 92, wherein said recombinant polypeptide is recombinant fibronectin fragment selected from the group consisting of CH-296 and

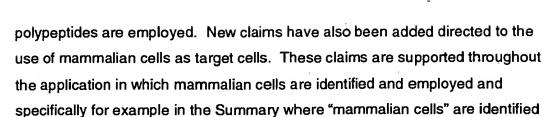
H-296

REMARKS

Entry of the above amendments is requested. Upon entry of the amendments, this application will contain claims 11-93. These amendments introduce no new subject matter to the application. The majority of the new claims are closely drawn from corresponding claims in the grandparent (PCT) application. New claims have been added directed to the recombinant polypeptide being selected from certain recombinant fibronectin polypeptides (e.g. CH-296, etc.). These claims are supported for example at page 16 of the application and in the specific Examples in which these recombinant

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Favorable consideration of the amended claims is solicited.

as targets for efficient retroviral infection in accordance with the invention.

Respectfully submitted

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